

<b><u>classifier</u></b>	<b><u>loci (out of 41,148 CBP peaks)</u></b>	<b><u>remaining enhancers</u></b>	<b><u>% of prev</u></b>	<b><u>filter</u></b>
All CBP Millipore KCl peaks	41148	41148		all CBP peaks (pre-filtered to remove 159 peaks with abnormally high enrichment)
CBP replicated by Ab2832	28004	27913	67.8	validated CBP peaks (2nd antibody)
TSS (RefSeq, Ensembl, UCSC,)	5079	25246	90.4	not an annotated TSS
longRNA	2438	24908	98.7	no initiation of a long (> 2kb) RNA (based on RNA-Seq)
spliced EST crossing TSS	3550	24752	99.4	not within 2000 bp of a 5' end of an EST that crosses a TSS
extragenic	18579	12899	52.1	at least 2kb from a gene
has H3K4me1 peak within 2kb	21378	5424	42.0	has H3K4me1 (& extragenic)
has H3K4me3 peak within 2kb (out of 41148)	6295	290	5.3	high H3K4me3 (& extragenic)
has H3K4me1 peak within 2kb	21378	7271		has H3K4me1 (& intragenic)